SAF-B03-018 ERDF Groundwater Well Samples FINAL DATA PACKAGE

MAIL COMPLETE COPY OF DATA PACKAGE TO:

Jim Rugg	H9-03	NB 12-6/05_ INITIAL/DATE
Rich Weiss	H9-01	NB 12-6/05_ INITIAL/DATE
Jeanette Duncan	Н9-02	NB 12-6/05_ INITIAL/DATE

COMMENTS:

SDG	K0047	SAF-B03-018
X Rad only	Chem only	Rad & Chem
X Complete	Par	tial





December 2, 2005

Ms. Joan Kessner Washington Closure Hanford 3190 George Washington Way MSIN H9-02 Richland, WA 99352

Reference:

P.O. #630

Eberline Services R5-10-085-7319, SDG K0047

Dear Ms. Kessner:

Enclosed is the data report for one water sample designated under SAF No. B03-018 received at Eberline Services on October 14, 2005. The sample was analyzed according to the accompanying chain-of-custody document.

Please call if you have any questions concerning this report.

Sincerely,

Melissa C. Mannion

Senior Program Manager

mu Marr

MCM/

Enclosure: Data Package

Analytical Services Tolf Free (800) 841-5487 www.eberlineservices.com

Case Narrative

Page 1 of 1

1.0 GENERAL

Washington Closure Hanford (WCH) Sample Delivery Group K0047 was composed of one water sample designated under SAF No. B03-018 with a Project Designation of: ERDF Groundwater Well Samples.

The sample was received as stated on the Chain-of-Custody document. Any discrepancies are noted on the Eberline Services Sample Receipt Checklist. The results were transmitted to WCH via e-mail on November 23, 2005.

2.0 ANALYSIS NOTES

2.1 Gross Alpha and Gross Beta Analysis

No problems were encountered during the course of the analyses.

2.2 Carbon-14 Analysis

No problems were encountered during the course of the analyses.

2.3 Iodine-129 Analysis

No problems were encountered during the course of the analyses.

2.4 Total Radium Analysis

No problems were encountered during the course of the analyses.

2.5 Technetium-99 Analysis

No problems were encountered during the course of the analyses.

2.6 Total Uranium Analysis

No problems were encountered during the course of the analyses.

Case Narrative Certification Statement

"I certify that this data package is in compliance with the SOW, both technically and for completeness, for other than the conditions detailed above. Release of the data obtained in this hard copy data package has been authorized by the Laboratory Manager or a designee, as verified by the following signature."

Melissa C. Mannion

Senior Program Manager

Mein Mann

/2/04/05-

E B E R L I N E S E R V I C E S / R I C H M O N D SAMPLE DELIVERY GROUP K0047

SDG <u>7319</u> Contact <u>Melissa C. Mannion</u> Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG_K0047</u>

SUMMARY DATA SECTION

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Prepared by

Meli Morm

Reviewed by

SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

REPORT GUIDE

Client	Hanford
Contract	No. 630
Case no	SDG_K0047

ABOUT THE DATA SUMMARY SECTION

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

SAMPLE SUMMARIES

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

WORK SUMMARY

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

METHOD BLANKS

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

LAB CONTROL SAMPLES

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

REPORT GUIDES

Page 1

SUMMARY DATA SECTION

Page 1

SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

GUIDE, cont.

Client	Hanford
Contract	No. 630
Case no	SDG K0047

ABOUT THE DATA SUMMARY SECTION

DUPLICATES

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

MATRIX SPIKES

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

DATA SHEETS

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

METHOD SUMMARIES

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

REPORT GUIDES

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

REPORT GUIDES
Page 2
SUMMARY DATA SECTION
Page 2

SAMPLE DELIVERY GROUP K0047

SDG	7319	
Contact	Melissa	C. Mannion

SAMPLE SUMMARY

Client	Hanford
Contract	No. 630
Case no	SDG_K0047

CLIENT SAMPLE ID	LOCATION	MATRIX LEVEL	LAB SAMPLE ID	SAF NO	CHAIN OF CUSTODY	COLLECTED
B1DMV8	DTS SAWS H93	WATER	R510085-01	B03-018	B03-018-124	10/13/05 09:54
Method Blank		WATER	R510085-03	B03-018		
Lab Control Sample		WATER	R510085-02	B03-018		
Duplicate (R510085-01)	DTS SAWS H93	WATER	R510085-04	B03-018		10/13/05 09:54
Spike (R510085-01)	DTS SAWS H93	WATER	R510085-05	B03-018		10/13/05 09:54

SAMPLE SUMMARY

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SUMMARY DATA SECTION

Page 3

SDG 7319
Contact Melissa C. Mannion

SAMPLE DELIVERY GROUP K0047

QC SUMMARY

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG K0047</u>

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	t SOLIDS	SAMPLE	BASIS AMOUNT	DAYS S		LAB SAMPLE ID	DEPARTMENT SAMPLE ID
7319	B03-018-124	BIDMV8	WATER		7.0 L		10/14/05	1	R510085-01	7319-001
		Method Blank Lab Control Sample Duplicate (R510085-01) Spike (R510085-01)	WATER WATER WATER WATER		7.0 L 7.0 L		10/14/05 10/14/05	1	R510085-03 R510085-02 R510085-04 R510085-05	7319-003 7319-002 7319-004 7319-005

OC SUMMARY
Page 1

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SUMMARY DATA SECTION

Page 4

SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

PREP BATCH SUMMARY

Client	Hanford
Contract	No. 630
Case no	SDG K0047

			PREPARATIO			- PLANCHETS ANALYZED					QUALI-	
TEST	MATRIX	METHOD	ватсн	20 %	CLIENT	MORE	RE	BLANK	LCS	DUP/ORIG	MS/ORIG	FIERS
Beta	Counting											
TC	WATER	Technetium 99 in Water	7148-075	10.0	1			1	1	1/1		
Gas I	Proportion	al Counting		<u>_</u>								
RAT	WATER	Total Alpha Radium in Water	7148-075	5.0	1			1	1	1/1		
Gas I	roportion	al Counting										
82B	WATER	Gross Beta in Water	7148-075	15.0	1		_	1	1	1/1		
88A	WATER	Gross Alpha in Water	7148-075	20.0	1			1	1	1/1	· · ·	
Gamma	Spectros	сору						<u> </u>				
I	WATER	Iodine 129 in Water	7148-075	5.0	1			1	1	1/1		
Kinet	ic Phosph	orimetry (KPA)										
U_T	WATER	Uranium, Total in Water	7148-075	9.0	1			1	1	1/1		
Liqui	d Scintil	lation Counting										
С	WATER	Carbon 14 in Water	7148-075	10.0	1			1	1	1/1	1/1	x

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.

Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

PREP BATCH SUMMARY
Page 1
SUMMARY DATA SECTION
Page 5

SAMPLE DELIVERY GROUP K0047

SDG 7319

Contact Melissa C. Mannion

WORK SUMMARY

Client Hanford
Contract No. 630
Case no SDG K0047

CLIENT SAMPLE : LOCATION CUSTODY	ID SAF No	MATRIX	LAB SAMPLE II COLLECTED RECEIVED	PLANCHET	Test	SUF-	ANAL VOED	REVIE WE D	P.V	MORNIOD
					IBD1	FIA	AMALIZED	KEVIEWED	BY	METHOD
B1DMV8			R510085-01	7319-001	82B/82		11/16/05	11/16/05	MWT	Gross Beta in Water
DTS SAWS H93		WATER	10/13/05	7319-001	88 A /88		11/18/05	11/23/05	MWT	Gross Alpha in Water
803-018-124	B03-018		10/14/05	7319-001	c		11/14/05	11/23/05	MWT	Carbon 14 in Water
				7319-001	I		11/18/05	11/22/05	MWT	Iodine 129 in Water
				7319-001	RAT		11/18/05	11/21/05	MWT	Total Alpha Radium in Water
				7319-001	TC		11/22/05	11/23/05	MWT	Technetium 99 in Water
				7319-001	ד_ט		11/07/05	11/09/05	MWT	Uranium, Total in Water
Method Blank			R510085-03	7319-003	82B/82		11/12/05	11/16/05	MWT	Gross Beta in Water
		WATER		7319-003	88A/88		11/18/05	11/23/05	MWT	Gross Alpha in Water
	B03-018			7319-003	С		11/14/05	11/23/05	MWT	Carbon 14 in Water
				7319-003	I		11/21/05	11/22/05	MWT	Iodine 129 in Water
				7319-003	RAT		11/18/05	11/21/05	MWT	Total Alpha Radium in Water
				7319-003	TC		11/21/05	11/23/05	MWT	Technetium 99 in Water
				7319-003	U_T		11/07/05	11/09/05	MWT	Uranium, Total in Water
ab Control Sam	nple		R510085-02	7319-002	82B/82		11/12/05	11/16/05	MWT	Gross Beta in Water
		WATER		7319-002	88A/88		11/22/05	11/23/05	MWT	Gross Alpha in Water
	B03-018			7319-002	С		11/14/05	11/23/05	MWT	Carbon 14 in Water
				7319-002	I		11/19/05	11/22/05	MWT	Iodine 129 in Water
				7319-002	RAT		11/18/05	11/21/05	MWT	Total Alpha Radium in Water
				7319-002	TC		11/21/05	11/23/05	MWT	Technetium 99 in Water
	·			7319-002	U_T		11/07/05	11/09/05	MWT	Uranium, Total in Water
uplicate (R510	085-01)		R510085-04	7319-004	82B/82		11/16/05	11/16/05	MWT	Gross Beta in Water
TS SAWS H93		WATER	10/13/05	7319-004	88A/88		11/18/05	11/23/05	MWT	Gross Alpha in Water
	B03-018		10/14/05	7319-004	С		11/14/05	11/23/05	MWT	Carbon 14 in Water
				7319-004	I		11/21/05	11/22/05	MWT	Iodine 129 in Water
				7319-004	RAT		11/18/05	11/21/05	MWT	Total Alpha Radium in Water
				7319-004	TC		11/22/05	11/23/05	MWT	Technetium 99 in Water
				7319-004	U_T		11/07/05	11/09/05	MWT	Uranium, Total in Water
pike (R510085-	01)		R510085-05	7319-005	С		11/14/05	11/23/05	MWT	Carbon 14 in Water
TS SAWS H93		WATER	10/13/05							
	B03-018		10/14/05							

WORK SUMMARY
Page 1
SUMMARY DATA SECTION
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 Lab id
 EBRLNE

 Protocol
 Hanford

 Version
 Ver 1.0

 Form
 DVD-CWS

 Version
 3.06

 Report date
 11/23/05

SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

WORK SUMMARY, cont.

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG K0047</u>

TEST	SAF No	COUNTS OF	TESTS BY SAM	CLIENT MORE	RE BLANK	LCS	DUP SPIKE	TOTAL
82B/82	B03-018	Gross Beta in Water	900.0_ALPHABETA_GPC	1	1	1	1	4
88A/88	B03-018	Gross Alpha in Water	900.0_ALPHABETA_GPC	1	1	1	1	4
С	B03-018	Carbon 14 in Water	C14_CHEM_LSC	1	1	1	1 1	5
I	B03-018	Iodine 129 in Water	I129_SEP_LEPS_GS	1	1	1	1	4
RAT	B03-018	Total Alpha Radium in Water	RATOT_GPC	1	ı	1	1	4
TC	B03-018	Technetium 99 in Water	TC99_TR_SEP_LSC	1	1	1	1	4
U_T	B03-018	Uranium, Total in Water	UTOT_KPA	1	1	1	1	4
TOTALS		100011111111111111111111111111111111111	0101_K14	7	7	7	7 1	

WORK SUMMARY
Page 2
SUMMARY DATA SECTION
Page 7

 Lab id
 EBRLNE

 Protocol
 Hanford

 Version
 Ver 1.0

 Form
 DVD-CWS

 Version
 3.06

 Report date
 11/23/05

EBERLINE SERVICES/RICHMOND SAMPLE DELIVERY GROUP K0047

R510085-03

METHOD BLANK

Method Blank

	7319 Melissa C. Mannion	Client/Case no Contract		SDG_K0047
Lab sample id Dept sample id		Client sample id Material/Matrix		STA WED
Dept sample id	7313-003	*	B03-018	WATER

ANALYTE	CAS NO	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	Test
Gross Alpha	12587-46-1	0.079	0.41	0.92	3.0	Ū	88A
Gross Beta	12587-47-2	-0.033	1.0	1.8	4.0	U	82B
Carbon 14	14762-75-5	-22.0	36	62	200	ប	С
Technetium 99	14133-76-7	-0.555	1.5	5.1	15	U	TC
Total Uranium (ug/L)	7440-61-1	0	0.023	0.054	0.10	U	UT
Total Radium	ALPHA-RA	-0.014	0.11	0.54	1.0	ប	RAT
Iodine 129	15046-84-1	0.805	1.8	4.0	5.0	U	I

ERDF Groundwater Well Samples

QC-BLANK	#54873

METHOD BLANKS
Page 1
SUMMARY DATA SECTION
Page 8

SAMPLE DELIVERY GROUP K0047

R510085-02

LAB CONTROL SAMPLE

Lab Control Sample

	7319 Melissa C. Mannion	Client/Case no Contract	Hanford No. 630	SDG_K0047
Lab sample id Dept sample id		Material/Matrix		WATER
		SAF No	B03-018	

ANALYTE	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST	ADDED pCi/L	2σ ERR pCi/L	REC	3σ LMTS (TOTAL)	PROTOCOL LIMITS
Gross Alpha	65.4	5.6	1.5	3.0		ABB	71.3	2.9	92	69-131	70-130
Gross Beta	65.0	3.3	1.9	4.0		82B	66.0	2.6	98	76-124	80-120
Carbon 14	7730	270	140	200		С	7970	320	97	83-117	80-120
Technetium 99	1150	30	5.3	15		TC	1090	44	106	83-117	80-120
Total Uranium (ug/L)	82.0	14	0.54	0.10		U_T	82.5	3.3	99	71-129	80-120
Total Radium	53.2	2.5	0.55	1.0		RAT	56.0	2.2	95	89-111	80-120
Todine 129	457	11	12	5.0		I	464	19	98	90-110	80-120

ERDF Groundwater Well Samples

QC-LCS #54872

LAB CONTROL SAMPLES
Page 1
SUMMARY DATA SECTION
Page 9

SAMPLE DELIVERY GROUP K0047

R510085-04

Dept sample id <u>7319-004</u>

DUPLICATE

B1DMV8

SDG 7319 Client/Case no <u>Hanford</u> Contact Melissa C. Mannion DUPLICATE ORIGINAL Lab sample id <u>R510085-04</u> Lab sample id <u>R510085-01</u>

Dept sample id <u>7319-001</u>

Received <u>10/14/05</u>

SDG K0047

Contract No. 630

Client sample id B1DMV8 Location/Matrix DTS SAWS H93 WATER

Collected/Volume 10/13/05 09:54 7.0 L Custody/SAF No <u>B03-018-124</u> <u>B03-018</u>

ANALYTE	DUPLICATE pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST	ORIGINAL pCi/L	2σ ERR (COUNT)	MDA pCi/L	QUALI- FIERS	RPD	3σ TOT	DER σ
Gross Alpha	0.675	0.40	0.51	3.0		88A	0.862	0.97	1.5	ט	24	209	0.3
Gross Beta	44.8	2.8	1.9	4.0		82B	44.6	2.9	2.0		0	35	
Carbon 14	3.59	35	58	200	U	c	19.9	36	60	υ	_		0.6
Technetium 99	70.8	6.0	5.6	15		TC	73.1	6.2	5.4		3	28	
Total Uranium (ug/L)	1.97	0.32	0.055	0.10		υт	2.00	0.33	0.055		2	40	0.1
Total Radium	-0.044	0.17	0.61	1.0	U	RAT	0.168	0.17	0.61	U	_	••	1.8
Iodine 129	2.89	1.9	4.3	5.0	Ū	ı	5.30	1.5	3.4	_	59	89	2.0

ERDF Groundwater Well Samples

QC-DUP#1 54874

DUPLICATES Page 1 SUMMARY DATA SECTION Page 10

Lab id EBRLNE Protocol <u>Hanford</u> Version <u>Ver 1.0</u> Form <u>DVD-DUP</u> Version 3.06

Report date <u>11/23/05</u>

SAMPLE DELIVERY GROUP K0047

R510085-05

MATRIX SPIKE

B1DMV8

SDG 7319

Contact Melissa C. Mannion

MATRIX SPIKE

Lab sample id <u>R510085-05</u>

Dept sample id <u>7319-005</u>

ORIGINAL

Lab sample id <u>R510085-01</u> Dept sample id <u>7319-001</u>

Received 10/14/05

Client/Case no <u>Hanford</u> <u>SDG K0047</u>

Contract No. 630

Client sample id <u>B1DMV8</u>

WATER Location/Matrix DTS SAWS H93

Collected/Volume <u>10/13/05 09:54 7.0 L</u> Custody/SAF No <u>B03-018-124</u> <u>B03-018</u>

ANALYTE	SPI KE pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST	ADDED pCi/L	2σ ERR pCi/L	ORIGINAL pCi/L	2σ ERR (COUNT)	REC 3σ LN	TS PROTOCOL L) LIMITS
Carbon 14	17000	580	250	200	х	С	23900	960	19.9	36	<u>71</u> 87-1	13 60-140

ERDF Groundwater Well Samples

QC-MS#1	54875		

MATRIX SPIKES Page 1 SUMMARY DATA SECTION Page 11

Lab id EBRLNE Protocol Hanford

Version <u>Ver 1.0</u>

Form DVD-MS Version 3.06

Report date <u>11/23/05</u>

EBERLINE SERVICES / RICHMOND SAMPLE DELIVERY GROUP K0047

R510085-01

DATA SHEET

B1DMV8

	1	7319 Melissa C. Mannion	Client/Case no Contract		SDG_K0047
i	Lab sample id Dept sample id Received		Client sample id Location/Matrix Collected/Volume Custody/SAF No	DTS SAWS H93 10/13/05 09:54	7.0 L B03-018

ANALYTE	CAS NO	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	Quali- Fiers	TEST
Gross Alpha	12587-46-1	0.862	0.97	1.5	3.0	U	88A
Gross Beta	12587-47-2	44.6	2.9	2.0	4.0		82B
Carbon 14	14762-75-5	19.9	36	60	200	ប	С
Technetium 99	14133-76-7	73.1	6.2	5.4	15		TC
Total Uranium (ug/L)	7440-61-1	2.00	0.33	0.055	0.10		U_T
Total Radium	ALPHA-RA	0.168	0.17	0.61	1.0	บ	RAT
Iodine 129	15046-84-1	5.30	1.5	3.4	5.0		I

ERDF Groundwater Well Samples

DATA SHEETS
Page 1
SUMMARY DATA SECTION
Page 12

SAMPLE DELIVERY GROUP K0047

Test	TC	Mat	rix	WATER
SDG	7319			
Contact	Melis	ssa	c.	Mannion

METHOD SUMMARY

TECHNETIUM 99 IN WATER
BETA COUNTING

Client	<u>Hanford</u>
Contract	No. 630
Contract	SDG K0047

RESULTS

	LAB	RAW SUF-	Technetium	
CLIENT SAMPLE ID	SAMPLE ID	TEST FIX PLANCHET	99	
Preparation batch 7148-	075			
B1DMV8	R510085-01	7319-001	73.1	
Method Blank	R510085-03	7319-003	σ	
Lab Control Sample	R510085-02	7319-002	ok	
Duplicate (R510085-01)	R510085-04	7319-004	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	raw Test		DA ALIQ /L L		DILU-	\$	EFF *		FWHM keV	 	PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-0)75 2 <i>g</i> pr	ep er	ror 10.0 %	Reference	Lab	Notebool	k #7148	, p	j. 075				·	
B1DMV8	R510085-01		5.	4 0.100			93		50		40	11/17/05	11/22	GRB-203
Method Blank	R510085-03		5.	0.100			99		50			11/17/05	11/21	GRB-203
Lab Control Sample	R510085-02		5.	3 0.100			97		50			11/17/05	11/21	GRB-202
Duplicate (R510085-01)	R510085-04		5.	6 0.100			97		50		40	11/17/05	11/22	GRB-201
Nominal values and limit	s from metho	кd	15	0.100			20-105		50		180			

PROCEDURES	REFERENCE	TC99_TR_SEP_LSC
1	CP-431	Technetium-99 Purification of Soil or Resin by
		Extraction Chromatography, rev 2
	CP-008	Heavy Element Electroplating, rev 9

METHOD SUMMARIES

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SUMMARY DATA SECTION

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 Lab id
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 Protocol
 Hanford

 Version
 Ver 1.0

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 DVD-CMS

 Version
 3.06

 Report date
 11/23/05

SAMPLE DELIVERY GROUP K0047

Test RAT Matrix WATER

SDG 7319

Contact Melissa C. Mannion

METHOD SUMMARY

TOTAL ALPHA RADIUM IN WATER
GAS PROPORTIONAL COUNTING

Client	Hanford
Contract	No, 630
Contract	SDG K0047

RESULTS

CLIENT SAMPLE ID	SAMPLE ID	RAW SUF- TEST FIX	PLANCHET	Total I	adium
Preparation batch 7148-	075		······································		
B1DMV8	R510085-01		7319-001	U	
Method Blank	R510085-03		7319-003	υ	
Lab Control Sample	R510085-02		7319-002	ok	
Duplicate (R510085-01)	R510085-04		7319-004	_	Ū

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE II	RAW D TEST	_	MDA pCi/L		PREP FAC		# YIELD	eff *		FWHM keV	DRIFT KeV		PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-0	75 2σ	prep eri	ror 5.	0 %	Reference	Lab i	Notebook	#7148	, pg	. 075						
B1DMV8	R510085-0	01		0.61	0.200			95		100			36	11/16/05	11/18	GAW-115
Method Blank	R510085-0	03		0.54	0.200			92		100				11/16/05	11/18	GAW-114
Lab Control Sample	R510085-0	02		0.55	0.200			92		100				11/16/05	11/18	GAW-114
Duplicate (R510085-01)	R510085-0	04		0.61	0.200			94		100			36	11/16/05	11/18	GAW-115
Nominal values and limits	from met	hod		1.0	0.200			20-105		100		-	180			

PROCEDURES	REFERENCE	RATOT_GPC
	DWP-880	Total Radium in Drinking Water, rev 0
L		_\

AVERAGES ± 2 SD MDA 0.58 ± 0.075

POR 4 SAMPLES YIELD 93 ± 3

METHOD SUMMARIES

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP K0047

Test 82B Matrix WATER

SDG 7319

Contact Melissa C. Mannion

METHOD SUMMARY

GROSS BETA IN WATER
GAS PROPORTIONAL COUNTING

Client <u>Hanford</u>

Contract <u>No. 630</u>

Contract <u>SDG K0047</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF		1: Gross Beta	2: Sum, Beta Emitters	RESULT RATIO (%) 2+1 2σ
Preparation batch 7148-	075					
B1DMV8	R510085-01	82	7319-001	44.6		
Method Blank	R510085-03	82	7319-003	ט		
Lab Control Sample	R510085-02	82	7319-002	ok		
Duplicate (R510085-01)	R510085-04	82	7319-004	ok		
Nominal values and limi	ts from metho	od Ri	OLs (pCi/L)	4.0		
ERDF Groundwater Well S	amples					Average

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST		OA ALIQ	PREP	DILU-	RESID mg	EFF				PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-	075 2ø pi	rep er	ror 15.0 %	Reference	Lab	Noteboo!	k #7148	, po	7. 075					
B1DMV8	R510085-01		2.0				81	-	100		34	11/10/05	11/16	GRB-105
Method Blank	R510085-03	82	1.8	0.300			61		100			11/10/05		
Lab Control Sample	R510085-02	82	1.9	0.300			61		100			11/10/05		
Duplicate (R510085-01)	R510085-04	82	1.9	0.300			79		100		34			
Nominal values and limit	s from metho	od	4.0	0.300			5-250		100		180		<u></u>	

PROCEDURES	REFERENCE	900.0_ALPHABETA_GPC
	CP-120	Gross Alpha and Gross Beta in Water, rev 6

AVERAGES ± 2 SD MDA 1.9 ± 0.16
FOR 4 SAMPLES RESIDUE 70 ± 22

METHOD SUMMARIES

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SAMPLE DELIVERY GROUP K0047

Test 88A Matrix WATER
SDG 7319
Contact Melissa C. Mannion

METHOD SUMMARY

GROSS ALPHA IN WATER
GAS PROPORTIONAL COUNTING

Client <u>Hanford</u>

Contract <u>No. 630</u>

Contract <u>SDG K0047</u>

RESULTS

CLIENT SAMPLE ID	SAMPLE ID	RAW SUF- TEST FIX	PLANCHET	Gross Alpha	
Preparation batch 7148-	075				
B1DMV8	R510085-01	88	7319-001	U	
Method Blank	R510085-03	88	7319-003	ū	
Lab Control Sample	R510085-02	88	7319-002	ok	
Duplicate (R510085-01)	R510085-04	88	7319-004	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	raw Test	SUF- FIX	MDA pCi/L		PREP FAC	DILU-	RESID mg	EFF %		FWHM keV	 	PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-	075 2σ pr	ep er	ror 20	.0 % F	Reference	Lab	Notebool	c #7146	, pg	j. 075					,
B1DMV8	R510085-01	88		1.5	0.300			78		100		36	11/10/05	11/18	GRB-114
Method Blank	R510085-03	88		0.92	0.300			59		100			11/10/05	11/18	GRB-116
Lab Control Sample	R510085-02	88		1.5	0.300			60		100			11/10/05	11/22	GRB-105
Duplicate (R510085-01)	R510085-04	88		0.51	0.300			80		400		36	11/10/05	11/18	GRB-216
Nominal values and limit	ts from metho	d		3.0	0.300			5-250		100		180			

PROCEDURES	REFERENCE	900.0_ALPHABETA_GPC
	CP-120	Gross Alpha and Gross Beta in Water, rev 6

AVERAGES ± 2 SD MDA 1.1 ± 0.97

FOR 4 SAMPLES RESIDUE 69 ± 23

METHOD SUMMARIES

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SAMPLE DELIVERY GROUP K0047

Test I Matrix WATER

SDG 7319

Contact Melissa C. Mannion

METHOD SUMMARY

IODINE 129 IN WATER
GAMMA SPECTROSCOPY

Client	Hanford
Contract	No. 630
Contract	SDG K0047

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX	PLANCHET	Iodine	129
Preparation batch 7148-	075				
B1DMV8	R510085-01		7319-001	5.30	
Method Blank	R510085-03		7319-003	υ	
Lab Control Sample	R510085-02		7319-002	ok	
Duplicate (R510085-01)	R510085-04		7319-004	ok	υ

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	_	MDA i/L		PREP FAC	DILU- TION	YIELD	EFF %		FWHM keV			PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-0	175 2σ pz	ep er	ror 5.0 %	Refe	rence L	ab N	Totebook	c #7148	, pg	j. 075		•				
B1DMV8	R510085-01		3	.4 0	. 250			73		1013			36	11/18/05	11/18	XSPEC-004
Method Blank	R510085-03		4	.0 0	. 250			74		536				11/18/05	11/21	XSPEC-004
Lab Control Sample	R510085-02		_12	°	. 250			61		230				11/18/05	11/19	XSPEC-004
Duplicate (R510085-01)	R510085-04		4	. 3 0	. 250			75		820			39	11/18/05	11/21	XSPEC-004
Nominal values and limit	s from metho	đ	5	.0 0	.250			20-105		300	100		180			

1	PROCEDURES	REFERENCE	I129_SEP_LEPS_GS
		CP-024	Iodine-129, Sample Dissolution, rev 5
		CP-530	Iodine-129 Purification, rev 1

AVERAGES ± 2 SD	MDA _	5.9_	±	8.1
FOR 4 SAMPLES	YIELD _	71	±	<u>13</u>

METHOD SUMMARIES

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP K0047

Test	<u>U T</u>	Matri	K WATER
SDG	7319		
ontact	Melis	sa C.	Mannion

METHOD SUMMARY

URANIUM, TOTAL IN WATER
KINETIC PHOSPHORIMETRY (KPA)

Client	Hanford
Contract	No. 630
Contract	SDG K0047

RESULTS

CLIENT SAMPLE ID	SAMPLE ID	RAW SUF- TEST FIX PLANCHET	Total Uranium	
Preparation batch 7148-07	75			···
B1DMV8	R510085-01	7319-001	2.00	
Method Blank	R510085-03	7319-003	σ	
Lab Control Sample	R510085-02	7319-002	ok	
Duplicate (R510085-01)	R510085-04	7319-00 4	ok	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	MDA ug/L	ALIQ L		DILU-	YIELD					PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-	075 2σ p	rep er	ror 9.6) % R	eference	Lab 1	Noteboo	k #7148	, p	g. 075					
B1DMV8	R510085-01			0.055	0.0200							25	11/07/05	11/07	KPA-001
Method Blank	R510085-03			0.054	0.0200								11/07/05	11/07	KPA-001
Lab Control Sample	R510085-02		_	0.54	0.0200								11/07/05	11/07	KPA-001
Duplicate (R510085-01)	R510085-04			0.055	0.0200							25	11/07/05	11/07	KPA-001
Nominal values and limit	ts from metho	od		0.10	0.0200						_	180			

PROCEDURES	REFERENCE	UTOT_KPA
	CP-044	Sample Preparation for Total Uranium by Kinetic
		Phosphorimetry, rev 6
	CP-929	Calibration of the Kinetic Phosphorimeter, rev 9
I .		

AVERAGES ± 2 SD MDA 0.18 ± 0.49

FOR 4 SAMPLES YIELD _____ ± _____

METHOD SUMMARIES

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP K0047

Test	<u>c</u>	Mat	ri	c <u>w</u>	ATER	
SDG	7319					
Contact	Melis	ssa	C.	Ma	nnion	_

METHOD SUMMARY

CARBON 14 IN WATER

LIQUID SCINTILLATION COUNTING

Client	Hanj			
Contract	No.	630		
Contract	SDG	K0047	····	

RESULTS

	LAB	RAW SUF-	-		
CLIENT SAMPLE ID	SAMPLE ID	TEST FIX	PLANCHET	Carbon	14
Preparation batch 7148-	075				
B1DMV8	R510085-01		7319-001	U	
Method Blank	R510085-03		7319-003	ט	
Lab Control Sample	R510085-02		7319-002	ok	
Duplicate (R510085-01)	R510085-04		7319-004	-	ט
Spike (R510085-01)	R510085-05		7319-005	LOW	х

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	raw Test	SUF- FIX po	MDA i/L	ALIQ L	PREP	DILU-	*	eff ¥		FWHM keV			PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7148-0)75 2σ pr	ep err	or 10.0	* Ref	erence	Lab	Notebool	c #7148	, pg	. 075						
B1DMV8	R510085-01		60		.0300			100		50			32	11/14/05	11/14	LSC-004
Method Blank	R510085-03		62	0	.0300			100		50				11/14/05	· •	
Lab Control Sample	R510085-02		140	0	.0300			100		10				11/14/05		
Duplicate (R510085-01)	R510085-04		58	0	.0300			100		50			32	11/14/05	11/14	LSC-004
Spike (R510085-01)	R510085-05		250	0	0200			100		7			32	11/14/05	11/14	LSC-004
Nominal values and limit	s from metho	d	200	0	. 0300	·				50		•	180			

PROCEDURES	REFERENCE	C14_CHEM_LSC
	CP-241	Carbon-14 in Aqueous Samples, rev 6

AVERAGES ± 2 SD	MDA	110	±	170
FOR 5 SAMPLES	YIELD	100	±	0

METHOD SUMMARIES

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SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

REPORT GUIDE

Client	Hanford
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SAMPLE SUMMARY

The Sample and QC Summary Reports show all samples, including QC samples, reported in one Sample Delivery Group (SDG).

The Sample Summary Report fully identifies client samples and gives the corresponding lab sample identification. The QC Summary Report shows at the sample level how the lab organized the samples into batches and generated QC samples. The Preparation Batch and Method Summary Reports show this at the analysis level.

The following notes apply to these reports:

- * LAB SAMPLE ID is the lab's primary identification for a sample.
- * DEPARTMENT SAMPLE ID is an alternate lab id, for example one assigned by a radiochemistry department in a lab.
- * CLIENT SAMPLE ID is the client's primary identification for a sample. It includes any sample preparation done by the client that is necessary to identify the sample.
- * QC BATCH is a lab assigned code that groups samples to be processed and QCed together. These samples should have similar matrices.

QC BATCH is not necessarily the same as SDG, which reflects samples received and reported together.

* All Lab Control Samples, Method Blanks, Duplicates and Matrix Spikes are shown that QC any of the samples. Due to possible reanalyses, not all results for all these QC samples may be relevant to the SDG. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.

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SAMPLE DELIVERY GROUP K0047

SDG 7319
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PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches in one Sample Delivery Group (SDG) with information necessary to check the completeness and consistency of the SDG.

The following notes apply to this report:

- * The preparation batches are shown in the same order as the Method Summary Reports are printed.
- * Only analyses of planchets relevant to the SDG are included.
- * Each preparation batch should have at least one Method Blank and LCS in it to validate client sample results.
- * The QUALIFIERS shown are all qualifiers other than U, J, B, L and H that occur on any analysis in the preparation batch. The Method Summary Report has these qualifiers on a per sample basis.

These qualifiers should be reviewed as follows:

- X Some data has been manually entered or modified. Transcription errors are possible.
- P One or more results are 'preliminary'. The data is not ready for final reporting.
- 2 There were two or more results for one analyte on one planchet imported at one time. The results in DVD may not be the same as on the raw data sheets.

Other lab defined qualifiers may occur. In general, these should be addressed in the SDG narrative.

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SAMPLE DELIVERY GROUP K0047

SDG <u>7319</u> Contact <u>Melissa</u> C. Mannion

REPORT GUIDE

Client	Hanford
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WORK SUMMARY

The Work Summary Report shows all samples, including QC samples, and all relevant analyses in one Sample Delivery Group (SDG). This report is often useful as supporting documentation for an invoice.

The following notes apply to this report:

- * TEST is a code for the method used to measure associated analytes. Results and related information for each analyte are on the Data Sheet Report. In special cases, a test code used in the summary data section is not the same as in associated raw data. In this case, both codes are shown on the Work Summary.
- * SUFFIX is the lab's code to distinguish multiple analyses (recounts, reworks, reanalyses) of a fraction of the sample. The suffix indicates which result is being reported. An empty suffix normally identifies the first attempt to analyze the sample.
- * The LAB SAMPLE ID, TEST and SUFFIX uniquely identify all supporting data for a result. The Method Summary Report for each TEST has method performance data, such as yield, for each lab sample id and suffix and procedures used in the method.
- * PLANCHET is an alternate lab identifier for work done for one test. It, combined with the TEST and SUFFIX, may be the best link to raw data.
- * For QC samples, only analyses that directly QC some regular sample are shown. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.
- * The SAS (Special Analytical Services) Number is a client or lab assigned code that reflects special processing for samples, such as rapid turn around. Counts of tests done are lists by SAS number since it is likely to affect prices.

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SAMPLE DELIVERY GROUP K0047

SDG 7319 Contact Melissa C. Mannion

REPORT GUIDE

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DATA SHEET

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- * TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- * The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- * ERRORs can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- * A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- * When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

U The RESULT is less than the MDA (Minimum Detectable Activity).

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SUMMARY DATA SECTION

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SAMPLE DELIVERY GROUP K0047

SDG 7319
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GUIDE, cont.

Client	Hani	ford	
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DATA SHEET

If the MDA is blank, the ERROR is used as the limit.

- J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.
- B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

- L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.
- H Similar to 'L' except the recovery was high.
- P The RESULT is 'preliminary'.
- X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.
- 2 There were two or more results available for this analyte. The reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

* An MDA is underlined if it is bigger than its RDL.

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GUIDE, cont.

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG K0047</u>

DATA SHEET

- * An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- * A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- * When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

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SAMPLE DELIVERY GROUP K0047

SDG 7319
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REPORT GUIDE

Client	Hani	ord	
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LAB CONTROL SAMPLE

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- * An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits for the recovery.
- * The recovery is underlined if it is outside either of these ranges.

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Lab id <u>EBRLNE</u>

Protocol <u>Hanford</u>

Version <u>Ver 1.0</u>

Form <u>DVD-RG</u>

Version <u>3.06</u>

Report date <u>11/23/05</u>

SAMPLE DELIVERY GROUP K0047

SDG <u>7319</u> Contact <u>Melissa C. Mannion</u>

REPORT GUIDE

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DUPLICATE

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTs divided by their average expressed as a percent.

If both RESULTs are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

* The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTs prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTs. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- * The second limit for the RPD is the larger of:
 - 1. A fixed percentage specified in the protocol.

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SAMPLE DELIVERY GROUP K0047

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GUIDE, cont.

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DUPLICATE

- 2. A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.
- * The RPD is underlined if it is greater than either limit.
- * If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

* The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit.

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SAMPLE DELIVERY GROUP K0047

SDG 7319
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REPORT GUIDE

Client	<u> Hanford</u>	
Contract	No. 630	
Case no	SDG_K0047	

MATRIX SPIKE

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.
 - If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.
- * An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.
 - An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.
- * REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The errors of the two RESULTs, including those introduced by rounding them prior to printing.
 - If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.
 - 2. The error of ADDED.
 - 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits

REPORT GUIDES

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SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

GUIDE, cont.

Client	Hani	ford
Contract	No.	630
Case no	SDG	K0047

MATRIX SPIKE

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

* The recovery is underlined (out of spec) if it is outside either of these ranges.

REPORT GUIDES

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SAMPLE DELIVERY GROUP K0047

SDG 7319
Contact Melissa C. Mannion

REPORT GUIDE

Client	Hanford
Contract	No. 630
Case no	SDG_K0047

METHOD SUMMARY

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

* Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

* The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

* If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- * Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- * Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

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SDG 7319
Contact Melissa C. Mannion

GUIDE, cont.

Client	Hanford
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Case no	SDG_K0047

METHOD SUMMARY

means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- * Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
- * If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.

MDAs are underlined if greater than the printed RDL.

- * Aliquots are underlined if less than the nominal value specified for the method.
- * Prepareation factors are underlined if greater than the nominal value specified for the method.
- * Dilution factors are underlined if greater than the nominal value specified for the method.
- * Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
- * Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
- * Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

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METHOD SUMMARY

- * Count times are underlined if less than the nominal value specified for the method.
- * Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- * Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- * Days Held are underlined if greater than the holding time specified in the protocol.
- * Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1÷3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

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METHOD SUMMARY

results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

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PNNL				·•		(CHAI	N OF	CUSTODY/	SAMPLE A	ANALYSIS I	REQUEST	}	B03-018-12	:4
Collector								Contact/Re	quester Krrk	17 (73/9	7	Telephone No.		FAX	\neg
SAF No. B03-018	BREWIN	lG}	OΛ	Sampling Origin (73/9)							Purchase Order/Charge Code				
Project Title ERDF Groundwa		. 1						ins	SAWS H	167		Ice Chest No. SHL	584 Temp.		\neg
Shipped To (Lab)) ics					· · · · · · · · · · · · · · · · · · ·	Method of				Bill of Lading/Air Bil	INO. 3-912 3-	+170500	\neg
<u>Eberline Services</u> Protocol									Prio	ority: 45 Days		Offsite Property No.	(((())	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	-1
GPP POSSIBLE SAMPI	LE HAZARI	S/RE	MARK	S				<u> </u>		SPECIAL INSTR	UCTIONS Hol	ld Time	Total Activity Exem	ption: Yes 🗹 No	
Sample No.	Lab ID	•	Da	te	T	ime	No/Type	Container			Sample Analysis			Preservative	
B1DMV8]	w	10-13	145	00	754	1x20-mi.	Р	Activity Scan					None	
B1DMV8		w				1	2x1000-n	nL G/P	Gross Alpha; Gross	Beta				HNO3 to pH	<2
B1DMV8	†	w			_		1x125-ml	- G/P	Carbon-14					None	
B1DMV8		w	<u> </u>		7		4x1000-n	nL G/P	lodine-129		 			None	
B1DMV8		w			1		1x1000-n	nL G/P	Total Radium					HNO3 to pH	<2
B1DMV8		W	1		7		1x250-m	- G/P	Technetium-99					HCl to pH <2	2
B1DMV8	<u> </u>	w		,	1		1x100-m	_ G/P	Total Uranium					HNO3 to pH	<2
			1		_	·									
		1						_ _ _							
			†												
															
			1						Received By					14	
Relinquished By 1 R. BREWI	Print	O.L.	Bru	Sign WK	to	<u>م</u>			FEI) E	Print	Sign	Date/Time	S = Soil	Matrix * DS = Drum	m Solid
Relinquished By	ED	モメ	C. Per	10	Γ.	105	Date	Time	Received By	10	14/05	Date/Time	SE = Sediment SO = Solid SL = Sludge	T = Tissu WI = Wine	e
Relinquished By			`			· I		Time .	Received By			Date/Time	W ≈ Water O = Oil A = Air	I. ≃ Lioui V = Vese X = Other	ctation
Relinquished By						 -	Date	/Time	Received By	<u></u>		Date/Time			
FINAL SAMPLE		Metho	d (e.g., R	eturn to	custo	mer, per	lab procedu	re, used in proc	<u> </u>	!	Disposed By		Dai	e/Time	



RICHMOND, CA LABORATORY SAMPLE RECEIPT CHECKLIST

Client: DNNL City MCHCKND State WA Date/Time received 0/4/65 0,55coc No. 803 - 018 - 124
Container I.D. No. SHL 584 Requested TAT (Days) 45 P.O. Received Yes [] No []
INSPECTION
1. Custody seals on shipping container intact? Yes [X] No [] N/A []
2. Custody seals on shipping container dated & signed? Yes [x] No [] N/A []
3. Custody seals on sample containers intact? Yes [y] No [] N/A []
4. Custody seals on sample containers dated & signed? Yes [\(\int \)] No [] N/A []
5. Packing material is: Wet [] Dry [🗸]
6. Number of samples in shipping container: Sample Matrix
7. Number of containers per sample: (Or see CoC)
8. Samples are in correct container Yes [Y] No []
9. Paperwork agrees with samples? Yes [X] No []
10. Samples have: Tape [] Hazard labels [] Rad labels [] Appropriate sample labels []/
11. Samples are: In good condition [] Leaking [] Broken Container [] Missing []
12. Samples are: Preserved [X] Not preserved [X] Preservative HN03, HEL
13. Describe any anomalies:
14. Was P.M. notified of any anomalies? Yes [] No [] Date
15. Inspected by Date: 0 1465 Time: 1790
Customer Customer Sample Sample No. cpm mR/hr Wipe No. cpm mR/hr wipe
Sample No. cpm mR/hr Wipe No. cpm mR/hr wipe
on Chamber Ser. No Calibration date
Alpha Meter Ser. No Calibration date
Deta/Gamma Meter Ser. No Calibration date